

APPROVED	D.G. FIG.
CLASS	SUBCLASS
DRAFTSMAN	

6007988

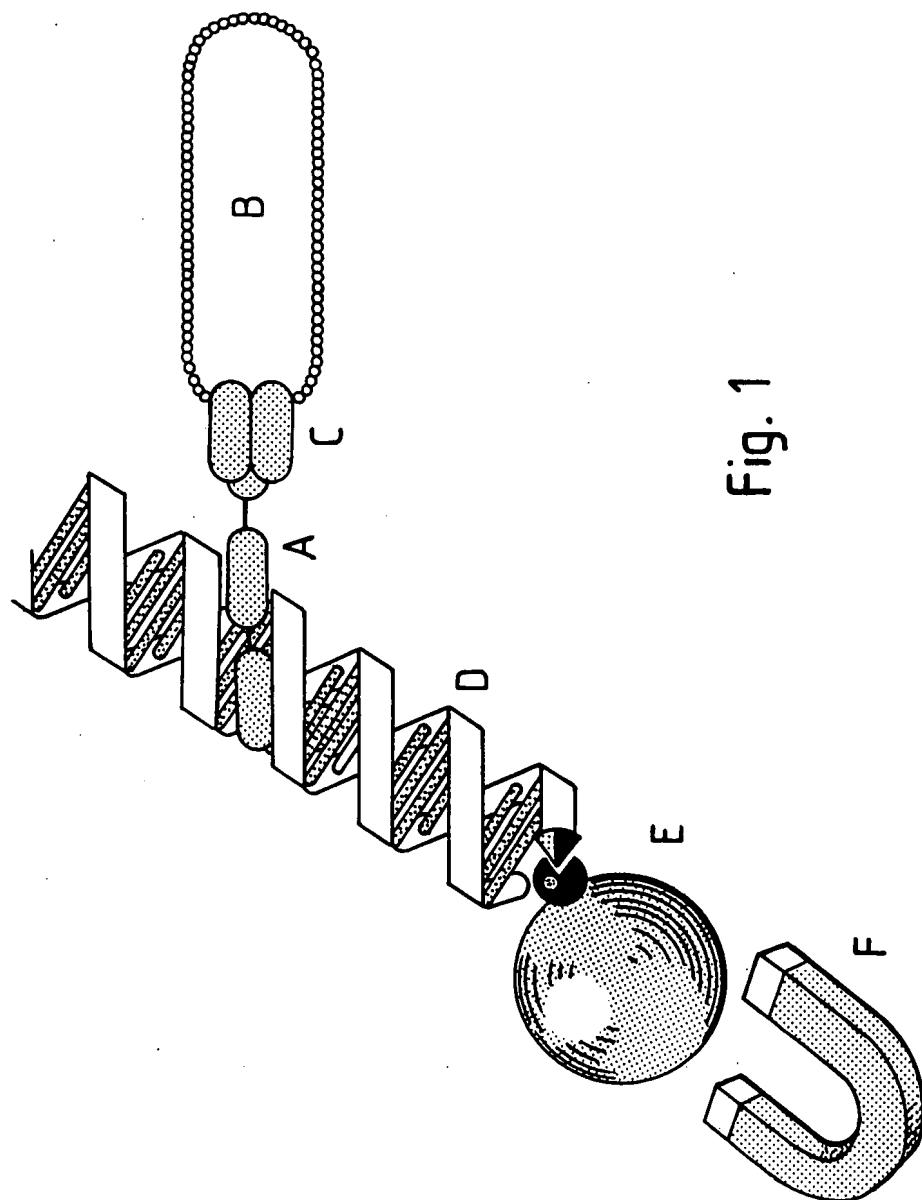


Fig. 1

APPROVED	O.G. FIG.	
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M A E E R P Θ A C P V E S G D R R F S R S D E L T R Θ T R I Θ T
 G Q K P F Q C R I - - G M R N F S X X X X L X X Θ X Θ K T Θ T
 G E K P F A C D I - - G G R K F A R S D E R K R Θ T K I Θ L R Q K D
 - 1 1 2 3 4 5 6 7 8 9

Fig. 2

APPROVED	O.G. FIG.	
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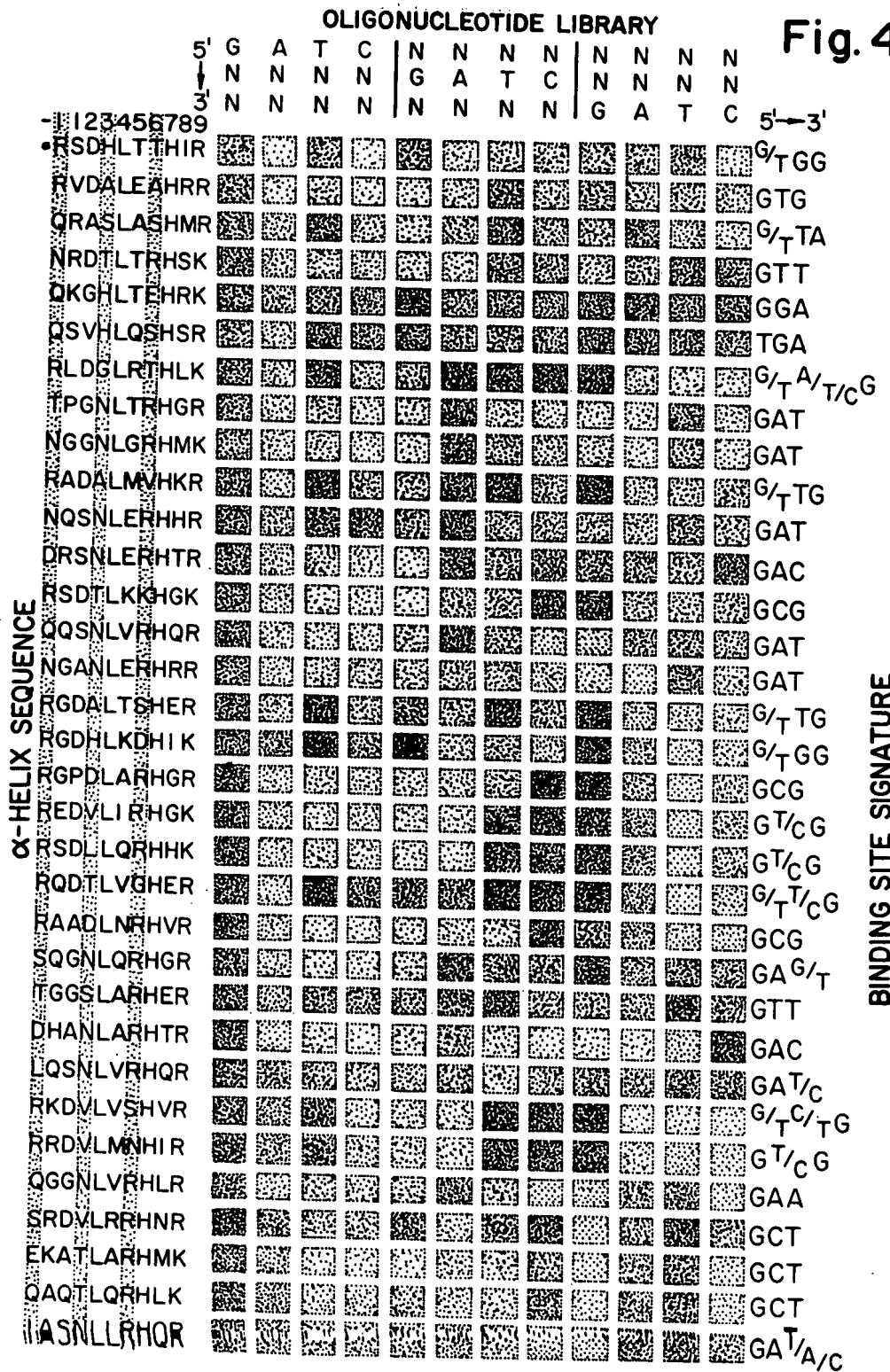
(i) TATGACTTGGATGGAGACCGCCTGG
ACTAACCTACCCCTCTGGCGGACCTTAA -

(ii) TATATA~~GGGGGG~~ATATATA
ATATAT~~GGGGGG~~ATATATGCG -

(iii) TATATAG~~GGGGGG~~ATATATA
ATATAT~~GGGGGG~~ATATATGCG -

Fig. 3

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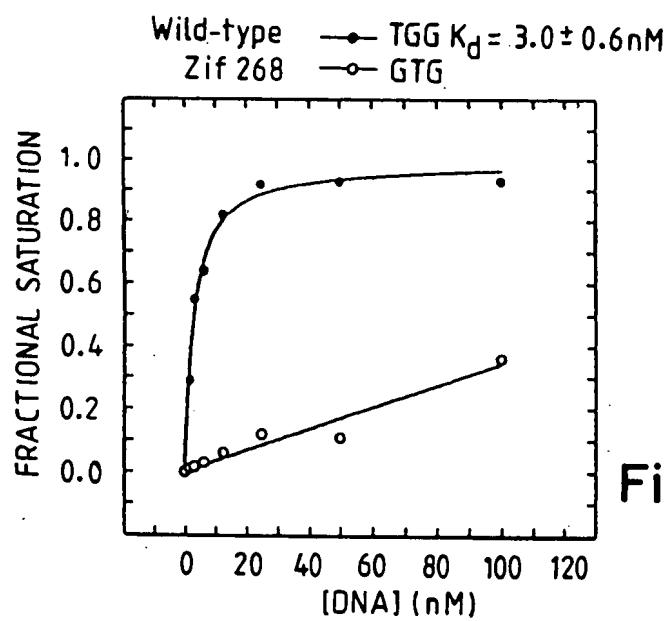


Fig. 5A

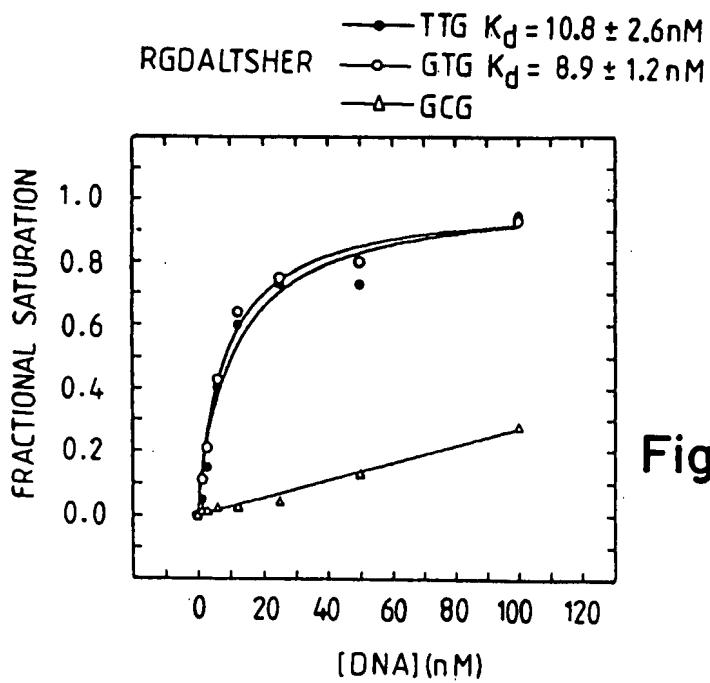


Fig. 5B

APPROVED	O.G. FIG.	
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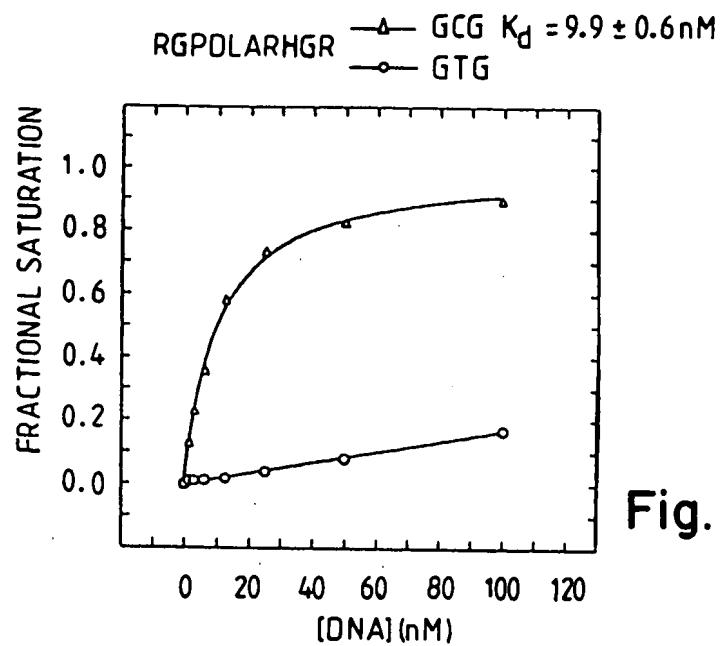


Fig. 5C

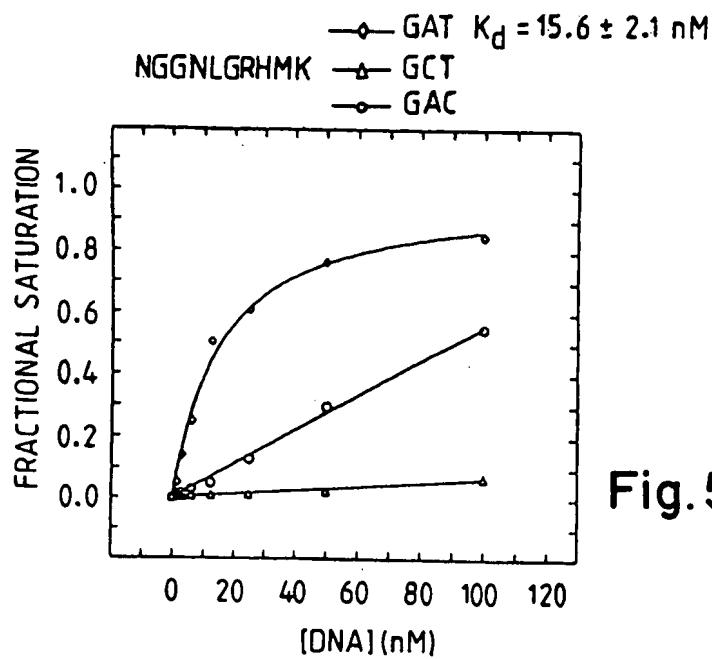


Fig. 5D

APPROVED	O.G. FIG.
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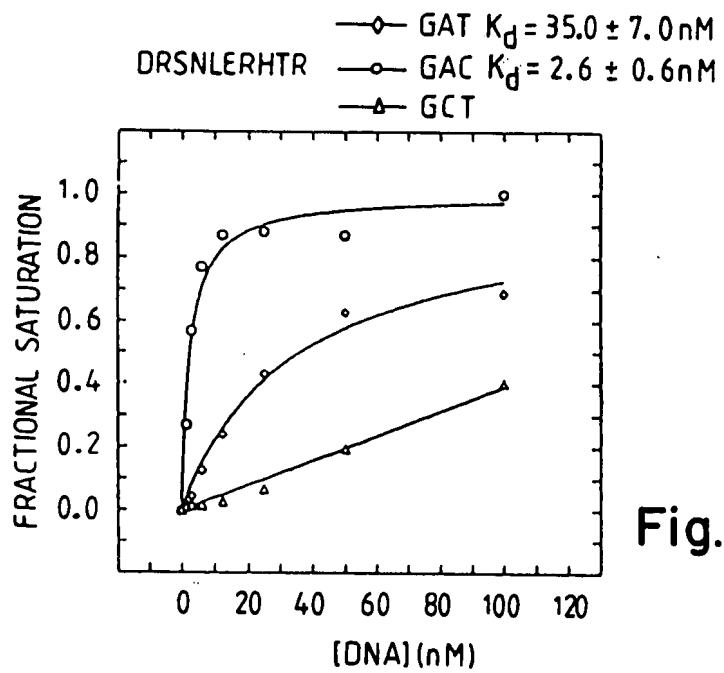


Fig. 5E

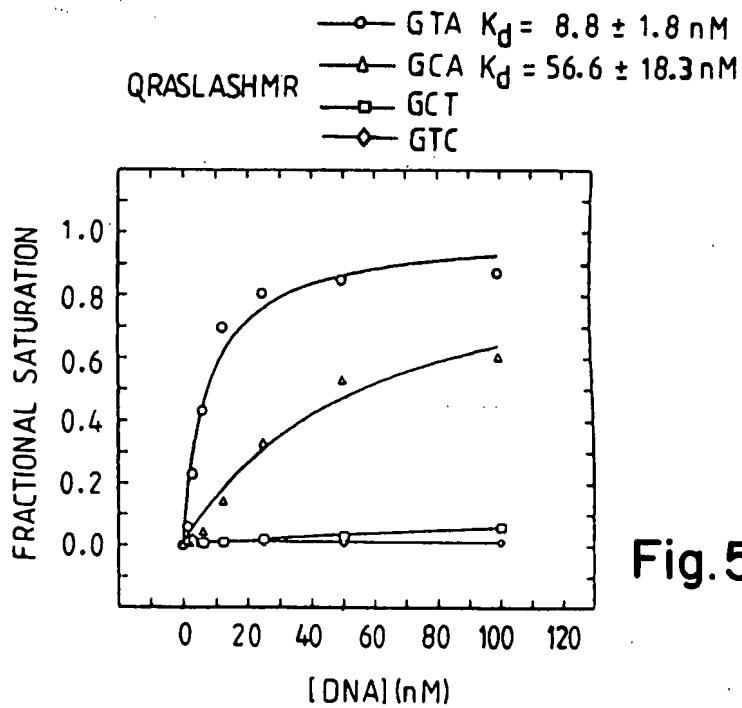


Fig. 5F

APPROVED	O.G. FIG.	
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1	<i>BCR - ABL</i>	TTC	CAT	GGA	GAC	GCA	<u>G</u>	<u>AA</u>	<u>GCC</u>	<u>CTT</u>	<u>CAG</u>	<u>CGG</u>	<u>CCA</u>
2	<i>BCR</i>	TTC	CAT	GGA	GAC	GCA	<u>G</u>	<u>gt</u>	<u>gag</u>	<u>tcc</u>	<u>ctc</u>	<u>acg</u>	<u>cca</u>
3	<i>ABL</i>	ccc	ctt	tct	cca	<u>g</u>	<u>AA</u>	<u>GCC</u>	<u>CTT</u>	<u>CAG</u>	<u>CGG</u>	<u>CCA</u>	

Fig. 6

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APPROVED	O.G. FIG.	
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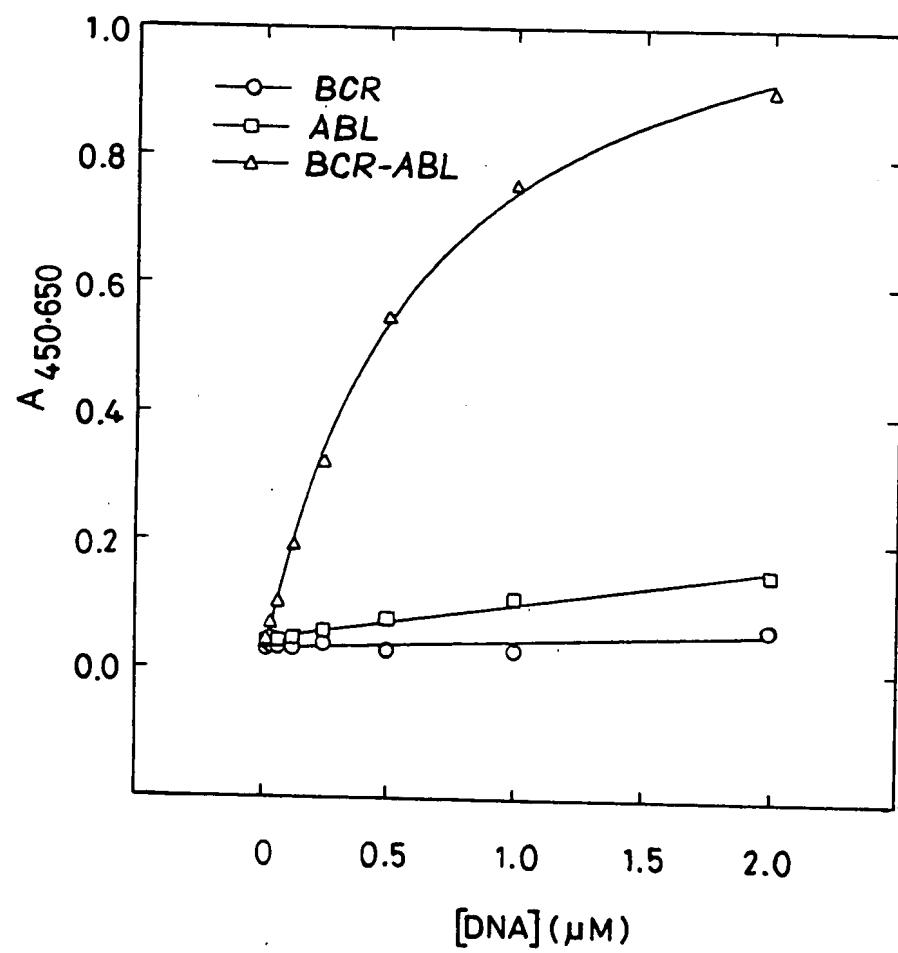


Fig. 8

APPROVED	O.G. FIG.	
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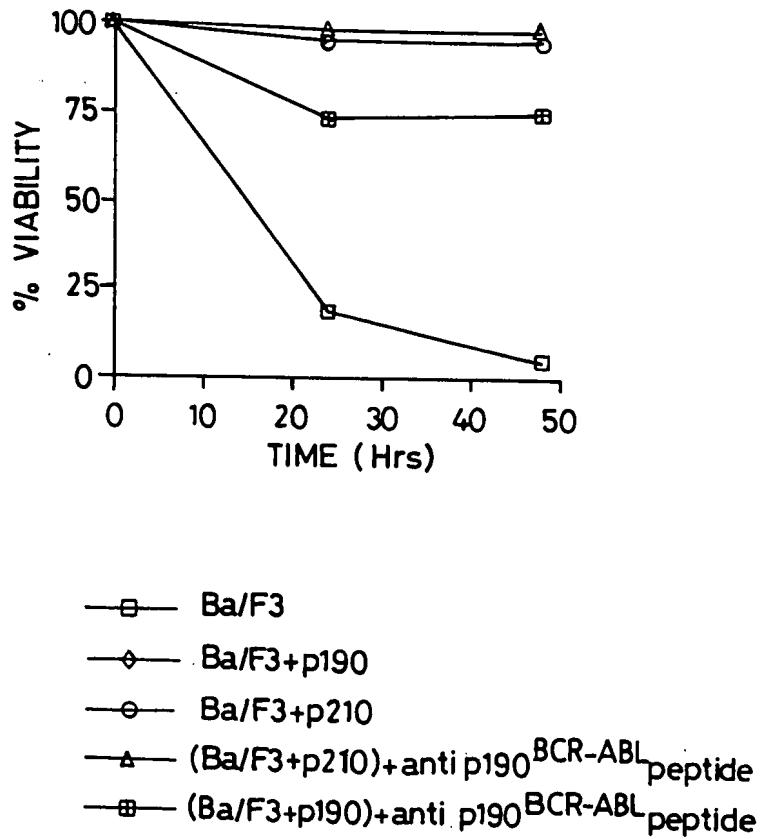


Fig. 9

APPROVED	O.G. FIG.	
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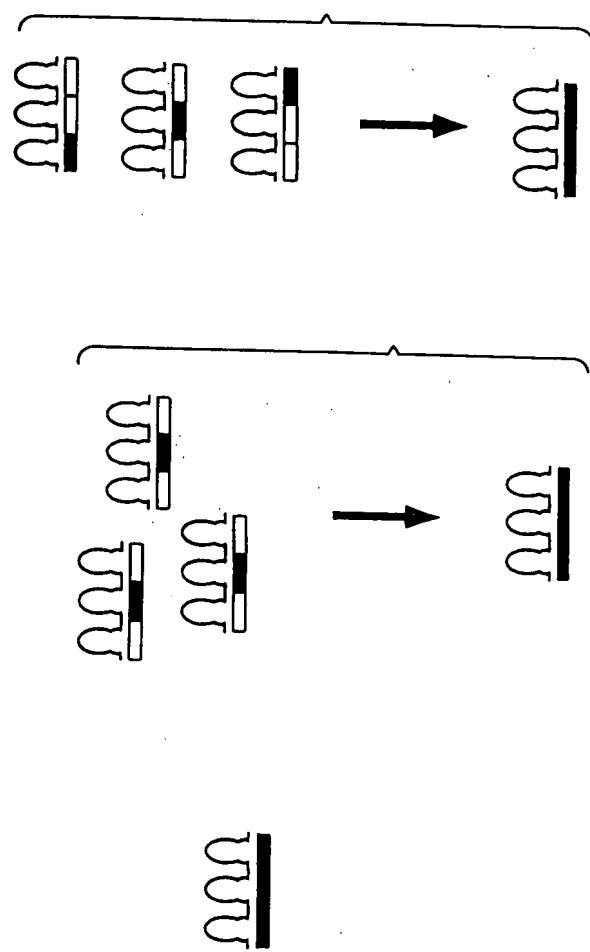


Fig. IOA

Fig. IOB

Fig. IOC

APPROVED	O.G. FIG.
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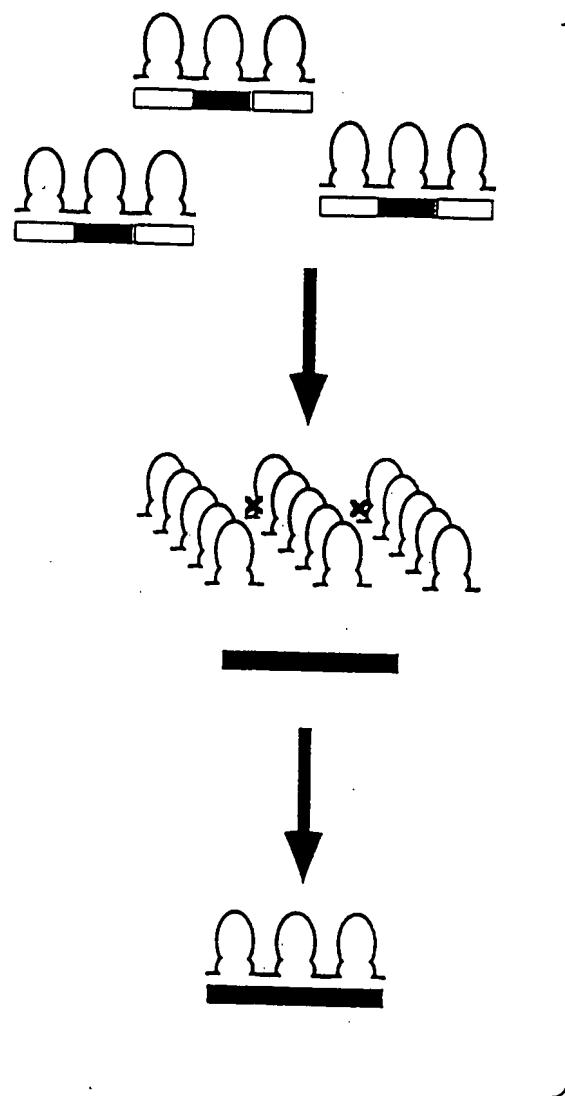


Fig.11

APPROVED	O.G. FIG.	
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Sequence logo for the C-terminal region of the protein. The y-axis shows positions -1, 1, 2, 3, 4, 5, 6, 7, 8, 9. The x-axis shows amino acids: M, A, E, E, K, P, F, Q, R, I, M, R, N, F, S. The logo is a 9x20 grid where height indicates frequency and shading indicates conservation. Arrows point to 'Cys' at position 1 and 'His' at position 9.

Fig. 2

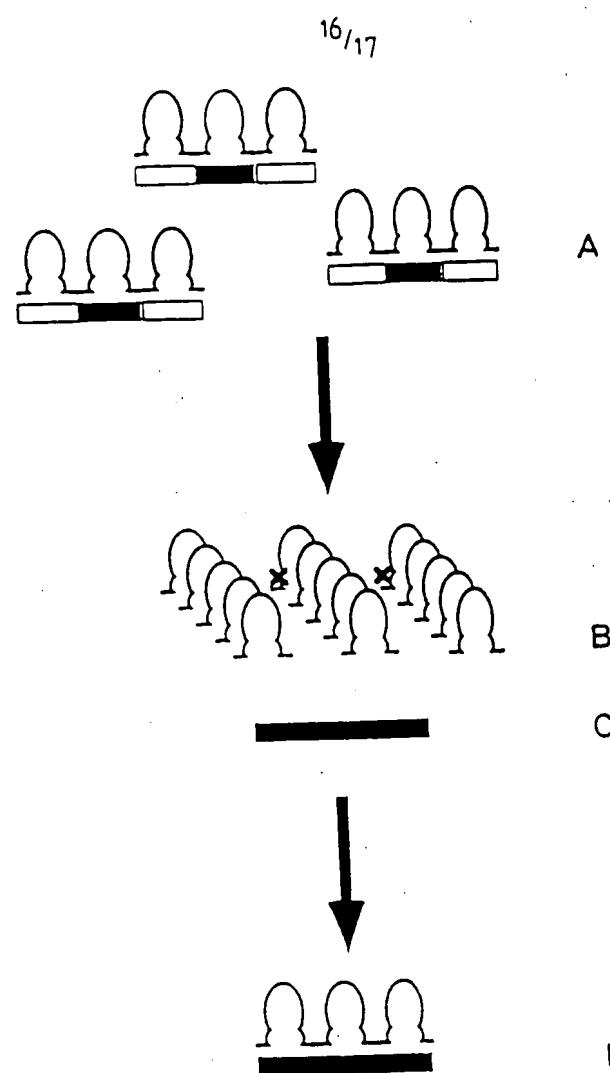


Fig. 14